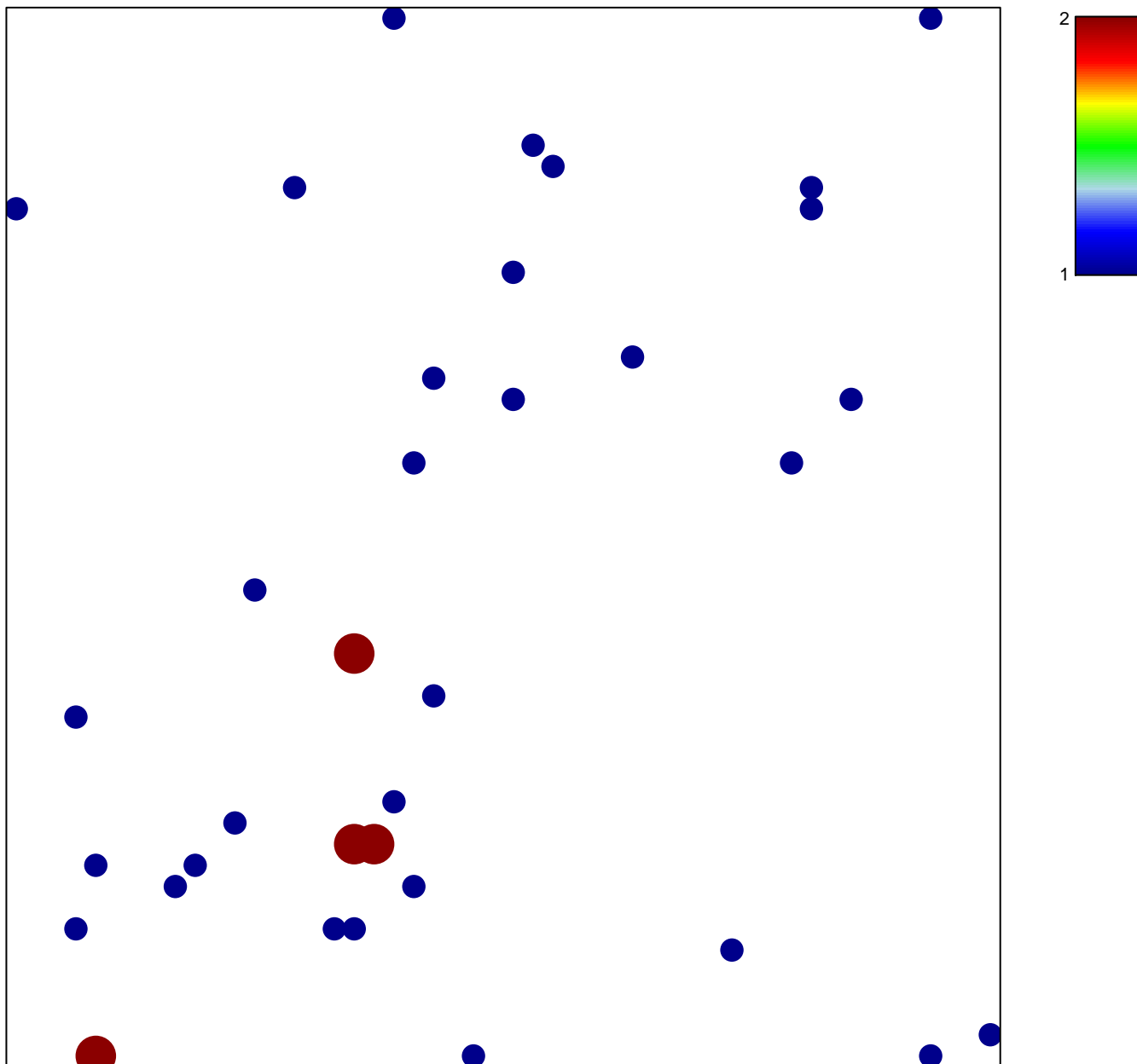


# SMAD protein signal transduction



# features = 39  
chi-square p = 0.82

# SMAD protein signal transduction



# features = 39 , max = 2